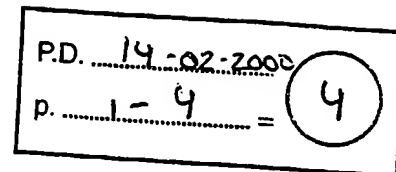


ID AC023078 standard; DNA; HTG; 164531 BP.  
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DT 14-FEB-2000 (Rel. 62, Created)  
DT 24-SEP-2001 (Rel. 69, Last updated, Version 8)  
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DE Homo sapiens chromosome 11 clone RP11-583F24 map 11, \*\*\* SEQUENCING IN  
DE PROGRESS \*\*\*, 2 ordered pieces.  
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KW HTG; HTGS\_ACTIVEFIN; HTGS\_FULLLTOP; HTGS\_PHASE2.  
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OS Homo sapiens (human)  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia;  
OC Eutheria; Primates; Catarrhini; Hominidae; Homo.  
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RA Birren B., Linton L., Nusbaum C., Lander E.;  
RT "Homo sapiens chromosome 11, clone RP11-583F24";  
RL Unpublished.  
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RN [2]  
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RA Birren B., Linton L., Nusbaum C., Lander E., Abraham H., Allen N.,  
RA Anderson S., Baldwin J., Barna N., Beckerly R., Beda F., Boguslavkiy L.,  
RA Boukhgalter B., Brown A., Burkett G., Castle A., Choepel Y., Colangelo M.,  
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RA Theodore J., Tirrell A., Vassiliev H., Viel R., Vo A., Wu X., Wyman D.,  
RA Ye W.J., Zimmer A., Zody M.;  
RT ;  
RL Submitted (08-FEB-2000) to the EMBL/GenBank/DBJ databases.  
RL Whitehead Institute/MIT Center for Genome Research, 320 Charles Street,  
RL Cambridge, MA 02141, USA  
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CC On Sep 20, 2001 this sequence version replaced gi:13559563.  
CC All repeats were identified using RepeatMasker:  
CC Smit, A.F.A. & Green, P. (1996-1997)  
CC <http://ftp.genome.washington.edu/RM/RepeatMasker.html>  
CC ----- Genome Center  
CC Center: Whitehead Institute/ MIT Center for Genome Research  
CC Center code: WIBR  
CC Web site: <http://www-seq.wi.mit.edu>  
CC Contact: [sequence\\_submissions@genome.wi.mit.edu](mailto:sequence_submissions@genome.wi.mit.edu)  
CC ----- Project Information  
CC Center project name: L6254  
CC Center clone name: 583\_F\_24  
CC -----  
CC \* NOTE: This is a 'working draft' sequence. It currently  
CC \* consists of 2 contigs. Gaps between the contigs  
CC \* are represented as runs of N. The order of the pieces  
CC \* is believed to be correct as given, however the sizes  
CC \* of the gaps between them are based on estimates that have  
CC \* provided by the submitter.  
CC \* This sequence will be replaced  
CC \* by the finished sequence as soon as it is available and  
CC \* the accession number will be preserved.  
CC \* 1 81101: contig of 81101 bp in length  
CC \* 81102 81201: gap of 100 bp  
CC \* 81202 164531: contig of 83330 bp in length.  
XX



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